

- 1 -

piece 1, NC_000913, ydbK_ydbJ+, config: linear, direction: +, begin: 1438779, end: 1439101

Sequence alignment of nine homologous genes from *S. enterica* serovar *Enteritidis*, *S. Typhimurium*, and *S. Infantis*. The genes are labeled *1438780 to *1438850. The alignment shows conserved amino acid residues in red, with a red dot at position 1438850. The sequence starts with a methionine (Met) residue.

-----1 NC 000913 vdbk

p35 4.3 bits

{ -----

|----- ... p35-p10 1438873 total 4.9

A screenshot of a web page from the Vivaldi browser. The page content includes the search term "ir ydbK_ydbJ+" highlighted in red. There are also green and blue annotations, likely from a tool like Gherkin Editor, pointing to specific parts of the text.

ir ydbK_ydbJ+

```
{-----} sd-(8)-ir 1438809 Gap 2.4 bits  
{-----} sd-ir 1438809 ydbK_ydbJ+ total 5.3 bits
```

[###> orf 19 codons

ns p35 3.8 bit

s p10 2.3 bits

... p35

```
... } p35-(24)-p10 1438873 Gap 2.4 bits  
... } p35-p10 1438873 total 4.9 bits
```

 p10 3.0 bits

{ --- ... p35-(23)-p10 1438962 Gap
--- ... p35-p10 1438962 total 4.2

{--- ... p35-(23)-p10 1438962 Gap
{--- p35-p10 1438962 total 4 2

```
{-----} p35-(23)-p10 1438901 Gap 1.4 bits  
|-----| p35-p10 1438901 total 4.6 bits
```

... p35 1.8 bits

p10 3.8 bits

... } p35-(23)-p10 1438962 Gap 1.4 bits
p35-p10 1438962 total 4.2 bits

The diagram illustrates the bacterial ribosome with various tRNA molecules (labeled tRNA₁ through tRNA₁₀) and their corresponding amino acids. The amino acid sequences are as follows:

- tRNA₁: *1439060
- tRNA₂: *1439070
- tRNA₃: *1439080
- tRNA₄: *1439090
- tRNA₅: *1439100
- tRNA₆: fMet - arg - ala - ala - phe - trp -
- tRNA₇: glu - asp - gln - arg - lys - lys - arg - cys - glu - gln - arg - phe - gly -
- tRNA₈: leu - lvs - ile - arg - cly - arg - lys - asp - ala - ser - ser - val - leu - gly -
- tRNA₉: sd
- tRNA₁₀: [- - - - - ... NC_000913.ydbJ

{-----} sd-(9)-ir 1439082 Gap 2.3 bits
sd-ir 1439082 vdbK vdbI+ total 9.4 bits